

GenCore version 6.3
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OM nucleic - nucleic search, using sw model

Run on: June 26, 2009, 03:11:25 ; Search time 31 Seconds
 (without alignments)
 28686.456 Million cell
 updates/sec

Title: US-09-300-482-225
 Perfect score: 316
 Sequence: 1
 gataaggtgcgcacactgag.....gaatggaggaggagtacttcga 316

Scoring table: IDENTITY_NUC
 Gapop 10.0 , Gapext 1.0

Searched: 2888648 seqs, 1391227531 residues

Total number of hits satisfying chosen parameters: 5777296

Minimum DB seq length: 0
 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
 Maximum Match 100%
 Listing first 45 summaries

Database : Pending_Patents_NA_New:*
 1:
 /ABSS/Data/CRF/ptodata/2/pna/PCT_NEW_COMB.seq:*
 2:
 /ABSS/Data/CRF/ptodata/2/pna/US08_NEW_COMB.seq:*
 3:
 /ABSS/Data/CRF/ptodata/2/pna/US09_NEW_COMB.seq:*
 4:
 /ABSS/Data/CRF/ptodata/2/pna/US10_NEW_COMB.seq:*
 5:
 /ABSS/Data/CRF/ptodata/2/pna/US11_NEW_COMB.seq:*
 6:
 /ABSS/Data/CRF/ptodata/2/pna/US12_NEW_COMB.seq:*
 7:
 /ABSS/Data/CRF/ptodata/2/pna/US12_NEW_COMB.seq1:*
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